SEQUENCE LISTING

<110> Lowell, George VanCott, Thomas Birx, Deborah

<120> PROTEIN AND PEPTIDE VACCINES FOR INDUCING MUCOSAL IMMUNITY

<130> 40646-20002.00

<140> US 09/214,701

<141> 1999-09-30

<150> PCT/US 97/12253

<151> 1997-07-10

<150> US 60/021,687

<151> 1996-07-10

<160> 18

<170> FastSEQ for Windows Version 4.0

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<211> 868

<212> PRT

<213> Virus HIV-1

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Ala Asn Leu Trp Val Thr Val Tyr Tyr Gly Val Pro Val Trp Lys Glu
35 40 45

Ala Thr Thr Leu Phe Cys Ala Ser Asp Ala Lys Ala Tyr Asp Thr 50 55 60

Glu Ala His Asn Val Trp Ala Thr His Ala Cys Val Pro Thr Asn Pro 65 70 75 80

Asn Pro Gln Glu Val Val Leu Glu Asn Val Thr Glu Asn Phe Asn Met 85 90 95

Trp Lys Asn Asn Met Val Glu Gln Met His Glu Asp Ile Ile Ser Leu 100 105 110

Trp Asp Gln Ser Leu Lys Pro Cys Val Lys Leu Thr Pro Leu Cys Val 115 120 125

Thr Leu Asn Cys Thr Asp Leu Asn Thr Asn Asn Thr Thr Asn Thr Thr 130 135 , 140

Glu Leu Ser Ile Ile Val Val Trp Glu Gln Arg Gly Lys Gly Glu Met 145 150 155 160

Arg Asn Cys Ser Phe Asn Ile Thr Thr Ser Ile Arg Asp Lys Val Gln
165 170 175

Arg Glu Tyr Ala Leu Phe Tyr Lys Leu Asp Val Glu Pro Ile Asp Asp 180 185

Asn Lys Asn Thr Thr Asn Asn Thr Lys Tyr Arg Leu Ile Asn Cys Asn 195 200 205 Thr Ser Val Ile Thr Gln Ala Cys Pro Lys Val Ser Phe Glu Pro Ile 215 Pro Ile His Tyr Cys Thr Pro Thr Gly Phe Ala Leu Leu Lys Cys Asn 235 230 Asp Lys Lys Phe Asn Gly Thr Gly Pro Cys Thr Asn Val Ser Thr Val 245 250 Gln Cys Thr His Gly Ile Arg Pro Val Val Ser Thr Gln Leu Leu 265 260 Asn Gly Ser Leu Ala Glu Glu Glu Val Val Ile Arg Ser Glu Asn Phe 280 Thr Asn Asn Ala Lys Thr Ile Ile Val Gln Leu Asn Val Ser Val Glu 295 300 Ile Asn Cys Thr Arg Pro Asn Asn His Thr Arg Lys Arg Val Thr Leu 310 315 Gly Pro Gly Arg Val Trp Tyr Thr Thr Gly Glu Ile Leu Gly Asn Ile 330 325 Arg Gln Ala His Cys Asn Ile Ser Arg Ala Gln Trp Asn Asn Thr Leu 345 Gln Gln Ile Ala Thr Thr Leu Arg Glu Gln Phe Gly Asn Lys Thr Ile 360 Ala Phe Asn Gln Ser Ser Gly Gly Asp Pro Glu Ile Val Met His Ser . 380 375 Phe Asn Cys Gly Glu Phe Phe Tyr Cys Asn Ser Thr Gln Leu Phe 390 395 Asn Ser Ala Trp Asn Val Thr Ser Asn Gly Thr Trp Ser Val Thr Arg 410 Lys Gln Lys Asp Thr Gly Asp Ile Ile Thr Leu Pro Cys Arg Ile Lys 420 425 Gln Ile Ile Asn Arg Trp Gln Val Val Gly Lys Ala Met Tyr Ala Leu 440 Pro Ile Lys Gly Leu Ile Arg Cys Ser Ser Asn Ile Thr Gly Leu Leu . 455 460 Leu Thr Arg Asp Gly Gly Gly Glu Asn Gln Thr Thr Glu Ile Phe Arg 475 470 Pro Gly Gly Gly Asp Met Arg Asp Asn Trp Arg Ser Glu Leu Tyr Lys 495 490 485 Tyr Lys Val Val Lys Ile Glu Pro Leu Gly Val Ala Pro Thr Lys Ala 505 500 Lys Arg Arg Val Val Gln Arg Glu Lys Arg Ala Val Gly Met Leu Gly 520 Ala Met Phe Leu Gly Phe Leu Gly Ala Ala Gly Ser Thr Met Gly Ala 540 535 Thr Ser Met Ala Leu Thr Val Gln Ala Arg Gln Leu Leu Ser Gly Ile 555 550 Val Gln Gln Gln Asn Asn Leu Leu Arg Ala Ile Lys Ala Gln Gln His 565 570 Leu Leu Gln Leu Thr Val Trp Gly Ile Lys Gln Leu Gln Ala Arg Ile 585 Leu Ala Val Glu Arg Tyr Leu Lys Asp Gln Gln Leu Leu Gly Phe Trp 600 Gly Cys Ser Gly Lys Leu Ile Cys Thr Thr Ala Val Pro Trp Asn Ala 615 Ser Trp Ser Asn Lys Thr Leu Asp Gln Ile Trp Asn Asn Met Thr Trp 635 630 Met Glu Trp Asp Arg Glu Ile Asp Asn Tyr Thr His Leu Ile Tyr Thr 655 650 645 Leu Ile Glu Glu Ser Gln Asn Gln Gln Glu Lys Asn Gln Gln Glu Leu

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665
Leu Gln Leu Asp Lys Trp Ala Ser Leu Trp Thr Trp Ser Asp Ile Thr
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Lys Trp Leu Trp Tyr Ile Lys Ile Phe Ile Met Ile Val Gly Gly Leu
                        695
Ile Gly Leu Arg Ile Val Phe Ala Val Leu Ser Ile Val Asn Arg Val
                   710
                                        715
Arg Gln Gly Tyr Ser Pro Leu Ser Phe Gln Thr Leu Leu Pro Asn Pro
                725
                                    730
Arg Gly Pro Asp Arg Pro Glu Gly Thr Glu Glu Gly Gly Glu Arg
            740
                                 745
                                                    750
Gly Arg Asp Gly Ser Thr Arg Leu Val His Gly Phe Leu Ala Leu Val
                            760
                                                 765
Trp Asp Asp Leu Arg Ser Leu Cys Leu Phe Ser Tyr His Arg Leu Arg
                        775
                                             780
Asp Leu Leu Leu Ile Val Ala Arg Ile Val Glu Leu Leu Gly Arg Arg
                    790
                                        795
Gly Trp Glu Val Leu Lys Tyr Trp Trp Asn Leu Leu Gln Tyr Trp Ser
                805
                                    810
                                                        815
Gln Glu Leu Lys Asn Ser Ala Val Ser Leu Val Asn Val Thr Ala Ile
            820
                                825
Ala Val Ala Glu Gly Thr Asp Arg Val Ile Glu Val Val Gln Arg Ile
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                                                845
Tyr Arg Ala Phe Leu His Ile Pro Arg Arg Ile Arg Gln Gly Phe Glu
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                        855
Arg Ala Leu Leu
865
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Asn Ala Asn Pro Asn Ala Asn Pro Asn Ala Asn Pro Asn Ala Asn Pro
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Asn Val Asp Pro
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gatecegggt gaetgaetga
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Gly Asn Val Gln Ala Ala Lys Asp Gly Gly Asn Thr Ala Ala Gly Arg
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Cys Tyr Gly Val Pro Val Ala Gln Thr Gln Thr Gly
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<211> 48
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<400> 14
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Gln Thr Gln Thr Gly Val Pro Val Ala Gln Thr Gln Thr Gly Val Pro
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                                25
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Cys Lys Tyr Asn Ala Thr Lys Ala
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                 5
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<212> DNA
<213> Artificial Sequence
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2019/050

<220>
<223> Synthetic DNA hydrophobic decapeptide coding sequence
<400> 17
ggtggttact gcttcgttgc tctgctgttc tgag

34

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<213> Artificial Sequence

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<223> Complementary chain of the hydrophobic decapeptide encoding sequence

<400> 18 tcgactcaga acagcagagc aacgaagcag taaccacc

38